Exhibit A

NCBI Sequence Viewer v2.0

NCBI

PubMed Nucleotide Protein Genome

Nucleotide

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Range: from 38997 to 39983   Show whole sequence   ☐ Reverse complemented strand   Fe

Links

☐ 1: BA000043. Reports Geobacillus kaust...[gi:56378377]

Features   **Sequence**

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DEFINITION      Geobacillus kaustophilus HTA426 DNA, complete genome.
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VERSION         BA000043.1  GI:56378377
KEYWORDS        .
SOURCE          Geobacillus kaustophilus HTA426
  ORGANISM      Geobacillus kaustophilus HTA426
                Bacteria; Firmicutes; Bacillales; Bacillaceae; Geobacillus.
REFERENCE       1
  AUTHORS       Takami,H., Takaki,Y., Chee,G.J., Nishi,S., Shimamura,S., Suzuki,H.,
                Matsui,S. and Uchiyama,I.
  TITLE         Thermoadaptation trait revealed by the genome sequence of
                thermophilic Geobacillus kaustophilus
  JOURNAL       (er) Nucleic Acids Res. 32 (21), 6292-6303 (2004)
  PUBMED        15576355
REFERENCE       2  (bases 1 to 987)
  AUTHORS       Takami,H., Takaki,Y. and Chee,G.
  TITLE         Direct Submission
  JOURNAL       Submitted (25-JUN-2003) Hideto Takami, Japan Marine Science and
                Technology Center, Microbial Genome Analysis Research Group; 2-15
                Natsushima-cho, Yokosuka, Kanagawa 237-0061, Japan
                (E-mail:takamih@jamstec.go.jp,
                URL:http://www.jamstec.go.jp/jamstec-e/bio/exbase.html,
                Tel:81-46-867-9643, Fax:81-46-867-9645)
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Aug 15 2006 13:27:38